



## ASH1L gene

ASH1 like histone lysine methyltransferase

### Normal Function

The *ASH1L* gene, also known as *KMT2H*, provides instructions for making an enzyme, called lysine-specific methyltransferase 2H, that is found in many organs and tissues of the body. Lysine-specific methyltransferase 2H functions as a histone methyltransferase. Histone methyltransferases are enzymes that modify proteins called histones. Histones are structural proteins that attach (bind) to DNA and give chromosomes their shape. By adding a molecule called a methyl group to histones (a process called methylation), histone methyltransferases control (regulate) the activity of certain genes. Lysine-specific methyltransferase 2H appears to turn on (activate) certain genes that are especially important for development of the brain.

### Health Conditions Related to Genetic Changes

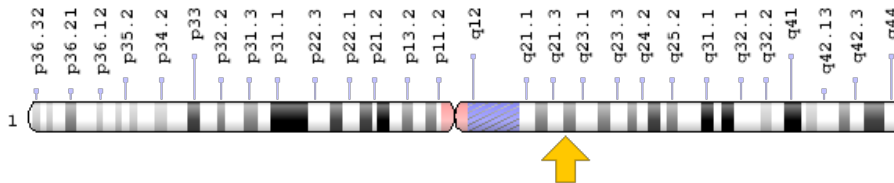
#### Autism spectrum disorder

At least seven *ASH1L* gene mutations have been identified in people with autism spectrum disorder (ASD), a varied condition characterized by impaired social skills, communication problems, and repetitive behaviors. Some *ASH1L* gene mutations associated with ASD change one building block (amino acid) in the lysine-specific methyltransferase 2H enzyme. Others delete genetic material in the *ASH1L* gene sequence or result in a premature stop signal that leads to an abnormally short enzyme. As a result of these mutations, the enzyme is nonfunctional. A lack of functional lysine-specific methyltransferase 2H enzyme disrupts histone methylation. The resulting changes in the expression of genes regulated by lysine-specific methyltransferase 2H affect brain development and increase the risk of ASD. Normal variations in other genes, as well as environmental risk factors, such as parental age, birth complications, and others that have not been identified, also affect an individual's risk of developing this complex condition.

## Chromosomal Location

Cytogenetic Location: 1q22, which is the long (q) arm of chromosome 1 at position 22

Molecular Location: base pairs 155,335,263 to 155,563,160 on chromosome 1 (Homo sapiens Updated Annotation Release 109.20200522, GRCh38.p13) (NCBI)



Credit: Genome Decoration Page/NCBI

## Other Names for This Gene

- absent small and homeotic disks protein 1 homolog
- ASH1
- ash1 (absent, small, or homeotic)-like
- ASH1-like protein
- ASH1L1
- histone-lysine N-methyltransferase ASH1L
- huASH1
- KMT2H
- lysine N-methyltransferase 2H
- probable histone-lysine N-methyltransferase ASH1L

## Additional Information & Resources

### Educational Resources

- Madame Curie Bioscience Database: Chromatin Mechanisms Regulating Gene Expression In Health And Disease  
<https://www.ncbi.nlm.nih.gov/books/NBK45032/>

### Scientific Articles on PubMed

- PubMed  
<https://www.ncbi.nlm.nih.gov/pubmed?term=%28%28ASH1L%5BTIAB%5D%29+OR+%28ASH1+like+histone+lysine+methyltransferase%5BTIAB%5D%29%29+OR+%28%28ASH1-like+protein%5BTIAB%5D%29+OR+%28ASH1%5BTIAB%5D%29+OR+%28absent+small+and+homeotic+disks+protein+1+homolog%5BTIAB%5D%29+OR+%28histone-lysine+N-methyltransferase+ASH1L%5BTIAB%5D%29+OR+%28huASH1%5BTIAB%5D%29+OR+%28lysine+N-methyltransferase+2H%5BTIAB%5D%29+OR+%28probable+histone-lysine+N-methyltransferase+ASH1L%5BTIAB%5D%29%29+AND+%28%28Genes%5BMH%5D%29+OR+%28Genetic+Phenomena%5BMH%5D%29%29+AND+english%5Bla%5D+AND+human%5Bmh%5D+AND+%22last+1800+days%22%5Bdp%5D>

### Catalog of Genes and Diseases from OMIM

- ASH1-LIKE HISTONE LYSINE METHYLTRANSFERASE  
<http://omim.org/entry/607999>

### Research Resources

- Atlas of Genetics and Cytogenetics in Oncology and Haematology  
[http://atlasgeneticsoncology.org/Genes/GC\\_ASH1L.html](http://atlasgeneticsoncology.org/Genes/GC_ASH1L.html)
- ClinVar  
<https://www.ncbi.nlm.nih.gov/clinvar?term=ASH1L%5Bgene%5D>
- HGNC Gene Symbol Report  
[https://www.genenames.org/data/gene-symbol-report/#!/hgnc\\_id/HGNC:19088](https://www.genenames.org/data/gene-symbol-report/#!/hgnc_id/HGNC:19088)
- Monarch Initiative  
<https://monarchinitiative.org/gene/NCBIGene:55870>
- NCBI Gene  
<https://www.ncbi.nlm.nih.gov/gene/55870>
- UniProt  
<https://www.uniprot.org/uniprot/Q9NR48>

### **Sources for This Summary**

- OMIM: ASH1-LIKE HISTONE LYSINE METHYLTRANSFERASE  
<http://omim.org/entry/607999>
- Eram MS, Kuznetsova E, Li F, Lima-Fernandes E, Kennedy S, Chau I, Arrowsmith CH, Schapira M, Vedadi M. Kinetic characterization of human histone H3 lysine 36 methyltransferases, ASH1L and SETD2. *Biochim Biophys Acta*. 2015 Sep;1850(9):1842-8. doi: 10.1016/j.bbagen.2015.05.013. Epub 2015 May 19.  
*Citation on PubMed:* <https://www.ncbi.nlm.nih.gov/pubmed/26002201>

- Rogawski DS, Ndoj J, Cho HJ, Maillard I, Grembecka J, Cierpicki T. Two Loops Undergoing Concerted Dynamics Regulate the Activity of the ASH1L Histone Methyltransferase. *Biochemistry*. 2015 Sep 8;54(35):5401-13. doi: 10.1021/acs.biochem.5b00697. Epub 2015 Aug 25.  
*Citation on PubMed:* <https://www.ncbi.nlm.nih.gov/pubmed/26292256>  
*Free article on PubMed Central:* <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC4664444/>
- Stessman HA, Xiong B, Coe BP, Wang T, Hoekzema K, Fenckova M, Kvarnung M, Gerdt J, Trinh S, Cosemans N, Vives L, Lin J, Turner TN, Santen G, Ruivenkamp C, Kriek M, van Haeringen A, Aten E, Friend K, Liebelt J, Barnett C, Haan E, Shaw M, Gecz J, Anderlid BM, Nordgren A, Lindstrand A, Schwartz C, Kooy RF, Vandeweyer G, Helsmoortel C, Romano C, Alberti A, Vinci M, Avola E, Giusto S, Courchesne E, Pramparo T, Pierce K, Nalabolu S, Amaral DG, Scheffer IE, Delatycki MB, Lockhart PJ, Hormozdiari F, Harich B, Castells-Nobau A, Xia K, Peeters H, Nordenskjöld M, Schenck A, Bernier RA, Eichler EE. Targeted sequencing identifies 91 neurodevelopmental-disorder risk genes with autism and developmental-disability biases. *Nat Genet*. 2017 Apr;49(4):515-526. doi: 10.1038/ng.3792. Epub 2017 Feb 13.  
*Citation on PubMed:* <https://www.ncbi.nlm.nih.gov/pubmed/28191889>  
*Free article on PubMed Central:* <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC5374041/>
- Wang T, Guo H, Xiong B, Stessman HA, Wu H, Coe BP, Turner TN, Liu Y, Zhao W, Hoekzema K, Vives L, Xia L, Tang M, Ou J, Chen B, Shen Y, Xun G, Long M, Lin J, Kronenberg ZN, Peng Y, Bai T, Li H, Ke X, Hu Z, Zhao J, Zou X, Xia K, Eichler EE. De novo genic mutations among a Chinese autism spectrum disorder cohort. *Nat Commun*. 2016 Nov 8;7:13316. doi: 10.1038/ncomms13316.  
*Citation on PubMed:* <https://www.ncbi.nlm.nih.gov/pubmed/27824329>  
*Free article on PubMed Central:* <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC5105161/>

---

Reprinted from Genetics Home Reference:  
<https://ghr.nlm.nih.gov/gene/ASH1L>

Reviewed: June 2017  
Published: June 23, 2020

Lister Hill National Center for Biomedical Communications  
U.S. National Library of Medicine  
National Institutes of Health  
Department of Health & Human Services